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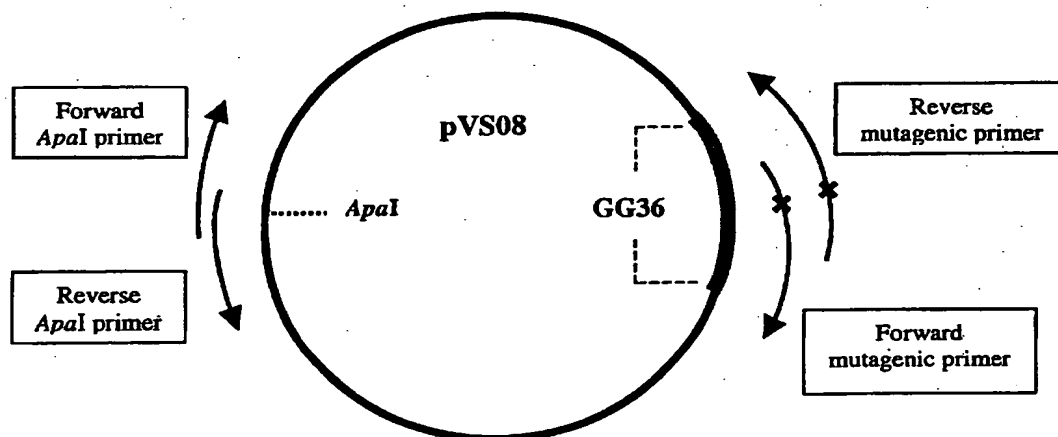
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(54) Title: MULTIPLY-SUBSTITUTED PROTEASE VARIANTS



(57) Abstract: Novel protease variants derived from the DNA sequences of naturally-occurring or recombinant non-human proteases are disclosed. The variant proteases, in general, are obtained by in vitro modification of a precursor DNA sequence encoding the naturally-occurring or recombinant protease to generate the substitution of a plurality of amino acid residues in the amino acid sequence of a precursor protease. Such variant proteases have properties which are different from those of the precursor protease, such as altered wash performance. The substituted amino acid residue equivalent to positions 7, 23, 26, 28, 29, 30, 31, 47, 66, 69, 73, 82, 85, 88, 90, 92, 93, 105, 113, 139, 148, 149, 150, 151, 178, 200, 201, 231, 233, 267 and/or 273 of *Bacillus amyloliquefaciens* subtilisin.

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